

**Nucleotides 1-1560 of the genome of
Enterovirus type 71 BrCr (GenBank Accession No.: U22521)**

1 TTTAAACAGC TGTGGGTTGT CACCCACCCA CAGGGTCCAC TGGGCGCTAG TACACTGGTA
61 TCTCGGTACC TTTGTACGCC TGTTTTATAC CCCCTCCCTG ATTTGCAACT TAGAAGCAAC

f1 (72-87)

121 GCAAACCAGA TCAATAGTAG GTGTGACATA CCAGTCCCAT CTTGATCAAG CACTTCTGTA

181 TCCCCGACACC GAGTATCAAT AGACTGTGCA CACGGTTGAA GGAGAAAACG TCCGTTACCC

f2 (167-187)

241 GGCTAATAC TTCGAGAAGC CTAGTAACGC CATTGAAGTT GCAGAGTGTT TCGCTCAGCA

f3 (248-266)

301 CTCCCCCGGT GTAGATCAGG TCGATGAGTC ACCGCATTCC CCACGGGCGA CCGTGGCGGT
361 GGCTGCGTTG GCGGCCTGCC TATGGGGTAA CCCATAGGAC GCTCTAATAC GGACATGGCG
421 TGAAGAGTCT ATTGAGCTAG TTAGTAGTCC TCCGCCCCCT GAATGCGGCT AATCCTAACT

f5 (423-439) p1 (448-474)

481 GCGGAGCACA TACCCTTAAT CCAAAGGGCA GTGTGTCGTA ACGGGCAACT CTGCAGCGGA

p2 (514-546)

541 ACCGACTACT TTGGGTGTCC GTGTTTCTTT TTATCTTGT ATTGGCTGCT TATGGTGACA

p3 (547-574) r2 (583-602)

601 ATTAAAGAAT TGTTACCATA TAGCTATTGG ATTGGCCATC CAGTGTCAAA CAGAGCTATT

r2 (583-602) r1 (627-645)

661 GTATATCTCT TTGTTGGATT CACACCTCTC ACTCTTGAAA CGTTACACAC CCTCAATTAC
721 ATTATACTGC TGAACACGAA GCGATGGGCT CCCAGGTCTC CACACAGCGA TCCGGCTCGC
781 ATGAGAATTC CAACTCAGCC ACGGAAGGCT CCACTATAAA TTACACAACC ATTAATTACT
841 ACAAAGACTC GTATGCTGCC ACTGCTGGAA AGCAAAGTCT CAAACAAGAT CCTGACAAGT
901 TTGCGAACCC TGTGAAGGAC ATCTTTACTG AAATGGCAGC GCCCTTAAAG TCTCCCTCTG
961 CTGAAGCATG TGGCTATAGC GACCGAGTGG CACAGCTTAC CATTGGAAAT TCCACCATTA
1021 CTACACAAGA AGCAGCAAAC ATAATAGTTG GGTATGGTGA GTGGCCTTCA TACTGCTCTG
1081 ATAATGATGC AACAGCGGTA GACAAACCTA CACGGCCTGA TGTCTCAGTA AATAGATTTT
1141 ACACGCTAGA CACTAAGCTA TGGGAGAAAT CATCCAAGGG GTGGTACTGG AAGTTCCCAG

f7 (1179-1198)

1201 ATGTACTGAC TGAAACCGGA GTTTTTGGTC CAAATGCACA ATTTCACTAC TTATACCGTT
1261 CAGGGTTCTG CATCCACGTT CAATGTAACG CTAGCAAATT TCACCAAGGG GCGCTACTCG
1321 TTGCGGTATT GCCCAGATAT GTCATTGGAA CAGTGGCAGG CGGCACAGGC ACAGAGAACA

16-1 (1343-1373)

1381 GTCACCCTCC TTATAAACAA ACCCAACCCG GCGCTGATGG ATTTGAATTA CAACATCCAT

71-2 (1390-1419)/16-2 (1390-1419)

1441 ATGTTCTTGA TGGTGGAAATT CCAATATCTC AGTTGACAGT GTGCCCTCAC CAGTGGATCA

71-3 (1453-1481) r4 (1485-1504)

1501 ATTTACGAAC CAACAATTGT GCCACCATAA TAGTGCCATA CATGAACACA CTACCTTTTG

Figure 1